

Go Clear Search Nucleotide for' Clipboard **Details** Limits History Preview/Index Show: 1 File default Get Subsequence Send to Display

1: AF232006[gi:8037790] This record was replaced or removed. See revision history for details.

20052 bp DNA linear BCT 23-MAY-2000 LOCUS AF232006

Pseudomonas syringae pv. tomato strain DC3000 AvrE (avrE), HrpW DEFINITION

(hrpW), and GstA (gstA) genes, complete cds; and unknown genes.

AF232006 ACCESSION

AF232006.1 GI:8037790 VERSION

KEYWORDS

Pseudomonas syringae pv. tomato SOURCE ORGANISM Pseudomonas syringae pv. tomato

Bacteria; Proteobacteria; gamma subdivision; Pseudomonadaceae;

Pseudomonas.

1 (bases 1 to 20052) REFERENCE

Charkowski, A.O., Alfano, J.R., Preston, G., Yuan, J., He, S.Y. and AUTHORS

Collmer, A.

The Pseudomonas syringae pv. tomato HrpW protein has domains TITLE

similar to harpins and pectate lyases and can elicit the plant

hypersensitive response and bind to pectate

J. Bacteriol. 180 (19), 5211-5217 (1998) JOURNAL

MEDLINE 98422476

(bases 1 to 20052) REFERENCE

Alfano, J.R., Charkowski, A.O., Deng, W.L., Badel, J.L., AUTHORS

Petnicki-Ocwieja, T., van Dijk, K. and Collmer, A.

The Pseudomonas syringae Hrp pathogenicity island has a tripartite TITLE

mosaic structure composed of a cluster of type III secretion genes

bounded by exchangeable effector and conserved effector loci that " |April 25,2000 -/

contribute to parasitic fitness and pathogenicity in plants

Proc. Natl. Acad. Sci. U.S.A. 97 (9), 4856-4861 (2000) JOURNAL

20243785 MEDLINE PUBMED 10781092

3 (bases 1 to 20052) REFERENCE

Alfano, J.R. and Collmer, A. AUTHORS

Direct Submission TITLE

Submitted (07-FEB-2000) Dept. Biol. Sci., UNLV, 1854 Maryland JOURNAL

Parkway, Las Vegas, NV 89154, USA

[WARNING] On Mar 14, 2001 this sequence was replaced by a newer COMMENT

version gi:13325077.

Location/Qualifiers FEATURES

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/organism="Pseudomonas syringae pv. tomato"

/strain="DC3000" /db xref="taxon:323"

/note="conserved effector locus flanking the hrp/hrc

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pathovar: tomato"

complement (1082..2569) CDS

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Revised: July 5, 2002.

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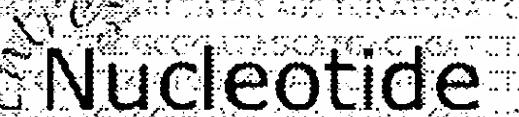
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PMC OMIM Taxonomy Nucleotide Protein Book **PubMed** Genome Structure Go Clear Search | Nucleotide for Limits Clipboard Details Preview/Index History Show: 20 **#File** default Send to Get Subsequence Display ☐ 1: AF232004. Pseudomonas syrin...[gi:13325077] Links 52498 bp DNA linear BCT 05-MAR-2001 LOCUS AF232004 Pseudomonas syringae pv. tomato strain DC3000 Hrp pathogenicity DEFINITION island, complete sequence. AF232004 AF061028 AF061029 AF232006 L41861 ACCESSION AF232004.3 GI:13325077 **VERSION** KEYWORDS Pseudomonas syringae pv. tomato SOURCE Pseudomonas syringae pv. tomato ORGANISM Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadaceae; Pseudomonas. 1 (bases 25494 to 29778) REFERENCE Preston, G., Huang, H.C., He, S.Y. and Collmer, A. AUTHORS The HrpZ proteins of Pseudomonas syringae pvs. syringae, glycinea, TITLE and tomato are encoded by an operon containing Yersinia ysc homologs and elicit the hypersensitive response in tomato but not soybean Mol. Plant Microbe Interact. 8 (5), 717-732 (1995) JOURNAL 96025089 MEDLINE 7579616 PUBMED 2 (bases 22134 to 25847; 29687 to 32670) REFERENCE Deng, W.L., Preston, G., Collmer, A., Chang, C.J. and Huang, H.C. AUTHORS Characterization of the hrpC and hrpRS operons of Pseudomonas TITLE syringae pathovars syringae, tomato, and glycinea and analysis of the ability of hrpF, hrpG, hrcC, hrpT, and hrpV mutants to elicit the hypersensitive response and disease in plants J. Bacteriol. 180 (17), 4523-4531 (1998) JOURNAL 98389667 MEDLINE PUBMED 9721291 3 (bases 31672 to 51723) REFERENCE Charkowski, A.O., Alfano, J.R., Preston, G., Yuan, J., He, S.Y. and AUTHORS Collmer, A. The Pseudomonas syringae pv. tomato HrpW protein has domains TITLE similar to harpins and pectate lyases and can elicit the plant hypersensitive response and bind to pectate J. Bacteriol. 180 (19), 5211-5217 (1998) JOURNAL MEDLINE 98422476 9748456 PUBMED REFERENCE 4 (bases 901 to 22404; 31672 to 51723) Alfano, J.R., Charkowski, A.O., Deng, W.L., Badel, J.L., AUTHORS Petnicki-Ocwieja, T., van Dijk, K. and Collmer, A. The Pseudomonas syringae Hrp pathogenicity island has a tripartite TITLE mosaic structure composed of a cluster of type III secretion genes bounded by exchangeable effector and conserved effector loci that 4/25/00 205. contribute to parasitic fitness and pathogenicity in plants Proc. Natl. Acad. Sci. U.S.A. 97 (9), 4856-4861 (2000)

5 (bases 1 to 52498) REFERENCE Ramos, A.R., Rehm, A.H. and Collmer, A.R. AUTHORS Pseudomonas syringae pv. tomato DC3000 hrpL through hrcU TITLE Unpublished JOURNAL 6 (bases 1 to 52498) REFERENCE Alfano, J.R. and Collmer, A. AUTHORS Direct Submission TITLE Submitted (07-FEB-2000) Dept. Biol. Sci., UNLV, 1854 Maryland JOURNAL Parkway, Las Vegas, NV 89154, USA 7 (bases 1 to 52498) REFERENCE Ramos, A.R., Rehm, A.H. and Collmer, A.R. AUTHORS Direct Submission TITLE Submitted (22-NOV-2000) Plant Pathology, Cornell University, 334 JOURNAL Plant Sciences Bldg., Ithaca, NY 14850, USA Sequence update by submitter REMARK 8 (bases 1 to 52498) REFERENCE Ramos, A.R., Rehm, A.H. and Collmer, A.R. AUTHORS Direct Submission TITLE Submitted (05-MAR-2001) Plant Pathology, Cornell University, 334 JOURNAL Plant Sciences Bldg., Ithaca, NY 14850, USA Sequence update by submitter REMARK On or before Mar 14, 2001 this sequence version replaced COMMENT gi:3228544, gi:790906, gi:3228541, gi:8037790, gi:11276506. Location/Qualifiers **FEATURES** 1..52498 source /organism="Pseudomonas syringae pv. tomato" /strain="DC3000" /db xref="taxon:323" /note="exchangeable effector locus flanking the hrp/hrc cluster pathovar: tomato" complement (908..1972) gene /gene="queA" complement (908..1972) CDS /gene="queA" /codon start=1 /transl table=11 /product="QueA" /protein id="AAF71483.1" /db xref="GI:8037772" /translation="MRVADFTFELPDSLIARHPLAERRSSRLLTLDGPTGALAHRQFT DLLEHLRSGDLMVFNNTRVIPARLFGQKASGGKLEILVERVLDSHRVLAHVRASKSPK PGSSILIDGGGEAEMVARHDALFELRFAEEVLPLLDRVGHMPLPPYIDRPDEGADRER YQTVYAQRAGAVAAPTAGLHFDQPLMEAIAAKGVETAFVTLHVGAGTFQPVRVEQIED HHMHSEWLEVSODVVDAVAACRARGGRVIAVGTTSVRSLESAARDGQLKPFSGDTDIF IYPGRPFHVVDALVTNFHLPESTLLMLVSAFAGYPETMAAYAAAIEHGYRFFSYGDAM FITRNPAPTAPQESAPEDHA" 2090..2177 tRNA /product="tRNA-Leu" 2343..2753 CDS /note="ORF4" /codon start=1 /transl table=11 /product="unknown" /protein id="AAF71484.1" /db xref="GI:8037773" /translation="MNKIVYVKAYFKPIGEEVSVKVPTGEIKKGFFGDKEIMKKETQW QQTGWSDCQIDGERLSKDVEDAVAQLNADGYEIQTVLPILSGAYDYALKYRYEIRHNR TELSPGDQSYVFGYGYSFTEGVTLVAKKFQSSAS" complement (2908..3324) CDS

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